SEQUENCE LISTING

<110> JONES, DAVID
      MANOS, ELIZABETH

<120>   TRDL-1 gamma, A NOVEL TUMOR NECROSIS-LIKE LIGAND

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<150>   60/157,913
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Trp
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Leu Ser Trp Gly Ala Ala Leu Gly Ala Val Ala Cys Ala Met Ala
Leu
          35   40   45

Leu Thr Gln Gln Thr Glu Leu Gln Ser Leu Arg Arg Glu Val Ser
Arg
     50   55   60

Leu Gln Gly Thr Gly Gly Pro Ser Gln Asn Gly Glu Gly Tyr Pro
Trp
65   70   75
80

Gln Ser Leu Pro Glu Gln Ser Ser Asp Ala Leu Glu Ala Trp Glu
Asn

                        85   90   95

Gly Glu Arg Ser Arg Lys Arg Arg Ala Val Leu Thr Gln Lys Gln
Lys
            100   105   110

Lys Gln His Ser Val Leu His Leu Val Pro Ile Asn Ala Thr Ser
Lys
            115   120   125

Asp Asp Ser Asp Val Thr Glu Val Met Trp Gln Pro Ala Leu Arg
Arg
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Ala
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Gly Val Tyr Leu Leu Tyr Ser Gln Val Leu Phe Gln Asp Val Thr
Phe
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Thr Met Gly Gln Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu
Thr
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Leu Phe Arg Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala
Tyr
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Asn Ser Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp
Ile
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Pro
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Trp
              20   25   30

Leu Ser Trp Gly Ala Ala Leu Gly Ala Val Ala Cys Ala Met Ala
Leu
            35   40   45

Leu Thr Gln Gln Thr Glu Leu Gln Ser Leu Arg Arg Glu Val Ser
Arg
          50   55   60

Leu Gln Gly Thr Gly Gly Pro Ser Gln Asn Gly Glu Gly Tyr Pro

Trp
65   70   75
80

Gln Ser Leu Pro Glu Gln Ser Ser Asp Ala Leu Glu Ala Trp Glu
Asn
              85   90   95

Gly Glu Arg Ser Arg Lys Arg Arg Ala Val Leu Thr Gln Lys Gln
Lys
              100   105   110

Lys Gln His Ser Val Leu His Leu Val Pro Ile Asn Ala Thr Ser
Lys
              115   120   125

Asp Asp Ser Asp Val Thr Glu Val Met Trp Gln Pro Ala Leu Arg
Arg
              130   135   140

Gly Arg Gly Leu Gln Ala Gln Gly Tyr Gly Val Arg Ile Gln Asp
Ala
145   150   155
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Gly Val Tyr Leu Leu Tyr Ser Gln Val Leu Phe Gln Asp Val Thr
Phe
              165   170   175

Thr Met Gly Gln Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu
Thr
              180   185   190

Leu Phe Arg Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala
Tyr
              195   200   205

Asn Ser Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp
Ile

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Leu Ser Val Ile Ile Pro Arg Ala Arg Ala Lys Leu Asn Leu Ser
Pro
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His Gly Thr Phe Leu Gly Phe Val Lys Leu
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Gly
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Asn Met Gly Gly Pro Val Arg Glu Pro Ala Leu Ser Val Ala Leu
Trp
                20   25   30

Leu Ser Trp Gly Ala Ala Leu Gly Ala Val Ala Cys Ala Met Ala
Leu
                35   40   45

Leu Thr Gln Gln Thr Glu Leu Gln Ser Leu Arg Arg Glu Val Ser
Arg
        50   55   60

Leu Gln Gly Thr Gly Gly Pro Ser Gln Asn Gly Glu Gly Tyr Pro
Trp
65   70   75
80

Gln Ser Leu Pro Glu Gln Ser Ser Asp Ala Leu Glu Ala Trp Glu
Asn
                85   90   95

Gly Glu Arg Ser Arg Lys Arg Arg Ala Val Leu Thr Gln Lys Gln
Lys
            100   105   110

Asn Asp Ser Asp Val Thr Glu Val Met Trp Gln Pro Ala Leu Arg
Arg
            115   120   125

Gly Arg Gly Leu Gln Ala Gln Gly Tyr Gly Val Arg Ile Gln Asp
Ala
    130   135   140

Gly Val Tyr Leu Leu Tyr Ser Gln Val Leu Phe Gln Asp Val Thr
Phe
145   150   155
160

Thr Met Gly Gln Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu
Thr
            165   170   175

Leu Phe Arg Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala
Tyr
            180   185   190

Asn Ser Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp
Ile
            195   200   205

Leu Ser Val Ile Ile Pro Arg Ala Arg Ala Lys Leu Asn Leu Ser
Pro
    210   215   220

His Gly Thr Phe Leu Gly Phe Val Lys Leu
225   230